

**AMENDMENTS TO THE SPECIFICATION**

Please replace the Sequence Listing filed March 18, 2010, with the substitute  
Sequence Listing submitted herewith.

Please amend Table 1 on page 7 of the specification filed May 7, 2007, as follows:

TABLE 1

Base Sequence of  $\mu$ -Chain Variable Region

GCTGAATTCTGGCTGACCAGGGCAGTCACCAGAGCTCCAGACAATGTCTG SEQ ID NO: 1  
TCTCCTTCCTCATCTTCCTGCCCCGTGCTGGGCCTCCCATGGGGTGTCTGT  
CACAGGTACAGCTGCAGCAGTCAGGTCCAGGACTGGTGAAGCCCGCGCAG  
ACCTCTCACTCACCTGTGCCATCTCCGGGGACAGTGTCTCTAGCAACAGT  
GCTACTTGGAACCTGGATCAGGCAGTCCCCATTGAGAGGCCTTGAGTGGCT  
GGGAAGGACATACTACAGGTCCAAGTGGTATAATGATTATGCAGTATCTG  
TGAAAAGTCGAATAACCATCAACCCAGACACATCCAAGAACCAGTTCTCCC  
TGCAGCTGAACTCTGTGACTCCCGAGGACACGGCTGTGTATTACTGTGCA  
AGAGAGAATTACTATGGTTCGGGGAGGTACAACTGGTTCGACCCCTGGGG  
CCAGGGAACCCTGGTCACCGTCTCCTCA

Base Sequence of  $\kappa$ -Chain Variable Region

TGTCAGGACACAGCATGGACATGAGGGTCCCCGCTCAGCTCCTGGGGCTC SEQ ID NO: 2  
CTGCTGCTCTGGTTCCCAGGTTCAGATGCGACATCCAGATGACCCAGTC  
TCCATCTTCCGTGTCTGCATCTGTAGGAGACAGAGTCACCATCACTTGTCC  
GGCGAGTCAGGGTATTAGCAGCTGGTTAGCCTGGTATCAGCAGAAACCAG  
GGAAAGCCCCTAAGCTCCTGATCTATGATGCATCCAGTTTGCAAAGTGGG  
GTCCCATCAAGGTTCAAGCGCAGTGGATCTGGGACAGATTTCACTCTCAC  
CATCAGCAGCCTGCAGCCTGAAGATTTTGCAACTTACTATTGTCAACAGGC  
TAACAGTTTCCCTCTCACTTTCGGCGGAGGGACCAAGGTGGAGATCAAA

Please amend Table 2 on pages 12-17 of the specification filed May 7, 2007, as follows:

TABLE 2: Examples of cDNA ~~encoding equivalent amino acids (i.e., SEQ ID NOs: 3 and 4 and conservatively modified variants thereof), encoding in the amino acid sequences~~ sequence of 9F11 antibody (SEQ ID NO: 7)

<u>SEQ ID NO: 7</u>	M S V S F L I F L P V L G L P W G V L S																			
	ATA TCT GTT TCT TTT TTA ATT TTT TTA CCT GTT TTA GGT TTA CCT TGA GGT GTT TTA TCT																			
<u>SEQ ID NO: 3</u>	ATG TCC GTC TCC TTC TTG ATC TTC TTG CCC GTC TTG GGC TTG CCC TGG GGC GTC TTG TCC																			
<u>SEQ ID NO: 4</u>	TCA GTA TCA	CTT	CTT CCA GTA CTT GGA CTT CCA	66A GTA CTT TCA																
	TCG GTG TCG	CTC	CTC CCG GTG CTC GGG CTC CCG	66G GTG CTC TCG																
	AGT AGT	CTA	CTA	CTA	CTA	CTA	CTA	CTA	CTA	CTA	CTA	CTA	CTA	CTA	CTA	CTA	CTA	CTA	AGT	CTA AGT
	AGC	AGC	CTG	CTG	CTG	CTG	CTG	CTG	CTG	CTG	CTG	CTG	CTG	CTG	CTG	CTG	CTG	CTG	AGC	CTG AGC

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Q V Q L Q Q S G P G L V K P A Q T L S L  
CAA GTT CAA TTA CAA CAA TCT GGT CCT GGT TTA GTT AAA CCT GCT CAA ACT TTA TCT TTA  
CAG GTC CAG TTG CAG CAG TCC GGC CCC GGC TTG GTC AAG CCC GGC CAG ACC TTG TCC TTG  
GTA CTT TCA GGA CCA GGA CTT GTA CCA GCA ACA CTT TCA CTT  
GTG CTC TCG GGG CCG GGG CTC GTG CCG GCG ACG CTC TCG CTC  
CTA AGT CTA CTA AGT CTA  
CTG AGC CTG CTG AGC CTG

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T C A I S G D S V S S N S A T W N W I R  
ACT TGT GCT ATT TCT GGT GAT TCT GTT TCT TCT AAT TCT GCT ACT TGA AAT TGA ATT CGT  
ACC TGC GCC ATC TCC GGC GAC TCC GTC TCC TCC AAC TCC GCC ACC TGG AAC TGG ATC CGC  
ACA GCA TCA GGA TCA GTA TCA TCA TCA GCA ACA CGA  
ACG GCG TCG GGG TCG GTG TCG TCG TCG GCG ACG CGG  
AGT AGT AGT AGT AGT  
AGC AGC AGC AGC AGC

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Q	S	P	L	R	G	L	E	W	L	G	R	T	Y	Y	R	S	K	W	Y
CAA	TCT	CCT	TTA	CGT	GGT	TTA	GAA	TGA	TTA	GGT	CGT	ACT	TAT	TAT	CGT	TCT	AAA	TGA	TAT
CAG	TCC	CCC	TTG	CGC	GGC	TTG	GAG	TGG	TTG	GGC	CGC	ACC	TAC	TAC	CGC	TCC	AAG	TGG	TAC
TCA	CCA	CTT	CGA	GGT	CTT	GGA	CGA	ACA	CTT	GGA	CGA	ACA	CTT	GGA	CGA	ACA	CTT	GGA	CGA
TCG	CCG	CTC	CGG	GGG	CTC	CTC	GGG	CGG	ACG	CTC	GGG	CGG	ACG	CTC	GGG	CGG	ACG	CTC	GGG
AGT	CTA	CTA	CTA	CTA	CTA	CTA	CTA	CTA	CTA	CTA	CTA	CTA	CTA	CTA	CTA	CTA	CTA	CTA	CTA
AGC	CTG	CTG	CTG	CTG	CTG	CTG	CTG	CTG	CTG	CTG	CTG	CTG	CTG	CTG	CTG	CTG	CTG	CTG	CTG

81

N D Y A V S V K S R I T I N P D T S K N  
AAT GAT TAT GCT GTT TCT GTT AAA TCT CGT ATT ACT ATT AAT CCT GAT ACT TCT AAA AAT  
AAC GAC TAC GCC GTC TCC GTC AAG TCC CGC ATC ACC ATC AAC CCC GAC ACC TCC AAG AAC  
GCA GTA TCA GTA TCA CGA ACA CCA ACA TCA  
GCG GTG TCG GTG TCG CGG ACG CCG ACG TCG  
AGT AGT AGT  
AGC AGC AGC

101

Q F S L Q L N S V T P E D T A V Y Y C A  
CAA TTT TCT TTA CAA TTA AAT TCT GTT ACT CCT GAA GAT ACT GCT GTT TAT TAT TGT GCT  
CAG TTC TCC TTG CAG TTG AAC TCC GTC ACC CCC GAG GAC ACC GCC GTC TAC TAC TGC GCC  
TCA CTT CTT TCA GTA ACA CCA ACA GCA GTA GCA  
TCG CTC CTC TCG GTG ACG CCG ACG GCG GTG GCG  
AGT CTA CTA AGT  
AGC CTG CTG AGC

121  
R E N Y Y G S G R Y N W F D P W G Q G T  
CGT GAA AAT TAT TAT GGT TCT GGT CGT TAT AAT TGA TTT GAT CCT TGA GGT CAA GGT ACT  
CGC GAG AAC TAC TAC GGC TCC GGC CGC TAC AAC TGG TTC GAC CCC TGG GGC CAG GGC ACC  
CGA GGA TCA GGA CGA CCA GGA GGA ACA  
CGG GGG TCG GGG CGG CCG GGG GGG ACG  
AGT  
AGC

141	L	V	T	V	S	S
	TTA	GTT	ACT	GTT	TCT	TCT
	TTG	GTC	ACC	GTC	TCC	TCC
	CTT	GTA	ACA	GTA	TCA	TCA
	CTC	GTG	ACG	GTG	TCG	TCG
	CTA				AGT	AGT
	CTG					AGC